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Fig. 1

E V Q L L E Q P G A  
GAGGTGCAGCTGCTCGAGCAGCCTGGGGCT 30

E L A K P G A S V K  
GAACTGGCAAAACCTGGGGCCTCAGTGAAG 60

M S C K A S G Y T F  
ATGTCCTGCAAGGCTTCTGGCTACACCTTT 90

T N Y W I H W V K Q  
ACTAACTACTGGATTCACTGGGTGAAACAG 120

R P G Q G L K W I G  
AGGCCTGGACAGGGTCTGAAATGGATTGGA 150

Y I N P A T G S T S  
TACATTAATCCTGCCACTGGTTCCACTTCT 180

Y N Q D F Q D R A T  
TACAATCAGGACTTTTCAGGACAGGGCCACT 210

L T A D K S S T T A  
TTGACCGCAGACAAGTCCTCCACCACAGCC 240

Y M Q L T S L T S E  
TACATGCAGCTGACCAGCCTGACATCTGAG 270

D S S V Y Y C A R E  
GACTCTTCAGTCTATTACTGTGCAAGAGAG 300

G Y D G F D S W G Q  
GGGTACGACGGGTTTGACTCCTGGGGCCAA 330

G T T L T V S S  
GGCACCACTCTCACAGTCTCCTCA 360

Fig. 2

E L V L T Q S P A I  
GAGCTCGTGCTCACCCAGTCTCCAGCAATC 30

M S A S P G E K V T  
ATGTCTGCATCTCCAGGGGAGAAGGTCACC 60

M T C S A S S S V N  
ATGACCTGCAGTGCCAGCTCAAGTGTAAT 90

Y M Y W Y Q Q K S G  
TACATGTACTTGGTACCAGCAGAAGTCAGGC 120

T S P K R W I Y D T  
ACCTCCCCCAAAGATGGATTTATGACACA 150

S K L A S G V P A R  
TCCAAATTGGCTTCTGGAGTCCCTGCTCGC 180

F S G S G S G T S Y  
TTCAGTGGCAGTGGGTCTGGGACCTCTTAC 210

S L T L S S M E A E  
TCTCTCACACTCAGCAGCATGGAGGCTGAA 240

D A A T Y Y C Q Q W  
GATGCCGCCACTTATTACTGCCCAGCAGTGG 270

S S N P Y T F G G G  
AGTAGTAATCCGTACACGTTCGGAGGGGGG 300

T K L E I K  
ACCAAGCTGGAGATAAAA 330

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Fig. 3

+1 E V Q L Q Q S G A E  
GAGGTTTCAGCTGCAGCAGTCTGGGGGCAGAG 30

+1 L V K P G A S V K L  
CTTGTGAAGCCTGGGGCCTCAGTCAAGTTG 60

+1 S C T S S G F N I K  
TCCTGCACATCTTCTGGCTTCAACATTAAA 90

+1 D T Y V H W M K Q R  
GACACCTATGTGCACTTGGATGAAACAGAGG 120

+1 P E Q G L E W I G K  
CCTGAACAGGGCCTGGAGTGGATTGGAAAG 150

+1 I D P A N G K T K Y  
ATTGATCCTGCGAATGGTAAAACTAAATAT 180

+1 D P I F Q A K A T M  
GACCCGATATTCCAGGCCAAGGCCACTATG 210

+1 T A D A S S N T A Y  
ACAGCAGACGCATCCTCCAATACAGCCTAC 240

+1 L Q L S S L T S E D  
CTGCAACTCAGCAGCCTGACTTCTGAGGAC 270

+1 T A V Y Y C A L P I  
ACTGCCGTCTATTACTGTGCTCTCCCCATT 300

+1 Y Y A S S W F A Y W  
TATTACGCTAGTTCCTGGTTTGCTTACTGG 330

+1 G Q G T L V T V S A  
GGCCAAGGGACTCTGGTCACTGTCTCTGCA 360

Fig. 4

+1 D I V M T Q S H K F  
GACATTGTGATGACCCAGTCTCACAAATTC 30

+1 M S T S V G D R V S  
ATGTCCACATCAGTAGGAGACAGGGGTCAGC 60

+1 I T C K A S Q D V G  
ATCACCTGCAAAGGCCAGTCAGGATGTGGGT 90

+1 T S V A W Y Q Q K P  
ACTTCTGTTGCCTGGTATCAACAGAAACCT 120

+1 G H S P K L L I Y W  
GGGCACTCTCCTAAATTACTGATTTACTTGG 150

+1 T S T R H T G V P D  
ACATCCACCCGGCACACTGGAGTCCCTGAT 180

+1 R F T G S G S G T D  
CGCTTCACAGGCAGTGGATCTGGGACAGAT 210

+1 F I L T I S N V Q S  
TTCATTCTCACCATTAGCAATGTGCAGTCT 240

+1 E D L A D Y F C Q Q  
GAAGACTTGGCAGATTATTTCTGTCAGCAA 270

+1 Y S S S P T F G G G  
TATAGCAGCTCTCCCACGTTTCGGAGGGGGG 300

+1 A K V E I K  
GCCAAGGTGGAAATAAAA 330

+1	D	I	L	L	T	Q	S	P	A	I	L	S	V	S	P	G	E	
	GACATCTTGC TGA																	50
+1	R	V	S	F	S	C	R	A	S	Q	S	I	G	T	R	I	H	
	AAGAGTCAGT TTCTCCTGGA																	100
+1	W	Y	Q	Q	R	T	N	G	S	P	R	L	L	I	K	Y		
	ACTGGTATCA ACAAAGAACA AATGGTTCTC CAAGGCTTCT CATAAAGTAT																	150
+1	G	S	E	S	I	S	G	I	P	S	R	F	S	G	S	G	S	
	GGTTCTGAGT CTATCTCTGG GATCCCTTCC AGGTTTAGTG GCAGTGGATC																	200
+1	G	T	D	F	S	L	S	I	N	S	V	E	S	E	D	I	A	
	AGGGACAGAT TTAGTCTTA GCATCAACAG TGTCGAGTCT GAAGATATTG																	250
+1	D	Y	Y	C	Q	Q	S	N	T	W	P	L	T	F	G	A		
	CAGATTATTA CTGTCAACAA AGTAATACCT GGCCGCTCAC GTTCGGTGCT																	300
+1	G	T	K	L	E	L	K											
	GGGACCAAGC TGGAGCTGAA A																	350

Fig. 5

+1	E	V	Q	L	L	E	E	S	G	G	G	L	V	K	P	G	G	
	GAGGTGCAGC TGCTCGAGGA GTCTGGGGGA GGCTTAGTGA AGCCTGGAGG 50																	
+1	S	L	Q	L	S	C	S	A	S	G	F	T	F	S	S	H	F	
	GTCCCTGCAA CTCTCCTGTT CAGCCTCTGG <u>ATTCACTTTC AGTAGCCATT</u> 100																	
+1	M	S	W	V	R	Q	T	P	E	K	R	L	E	W	V	A		
	<u>TCATGTCT</u> TG GGTTCGCCAA ACTCCAGAGA AGAGGCTGGA GTGGTTCGCA 150																	
+1	S	I	S	S	G	G	D	S	F	Y	P	D	S	L	K	G	R	
	<u>TCCATTAGTA GTGGTGGTGA CAGTTTCTAT CCAGACAGTC TGAAGGGC</u> CG 200																	
+1	F	A	I	S	R	D	N	A	R	N	I	L	F	L	Q	M	S	
	ATTCGCCATC TCCAGAGATA ATGCCAGGAA CATCCTGTTC CTGCAAATGA 250																	
+1	S	L	R	S	E	D	S	A	M	Y	F	C	T	R	D	Y		
	GCAGTCTGAG GTCTGAGGAC TCGGCCATGT ATTTCTGTAC AAGA <u>GACTAC</u> 300																	
+1	S	W	Y	A	L	D	Y	W	G	Q	G	T	S	V	T	V	S	
	<u>TCTTGGTATG CTTTGGACTA C</u> TGGGGTCAA GGAACCTCAG TCACCGTCTC 350																	
+1	S																	
	CTCA 400																	

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+1 E L Q M T Q S P S L S A S L G D  
GAGCTCCAGA TGACCCAGTC TCCATCCAGT CTGTCTGCAAT CCCTTGGAGA 50  
+1 T I T I T C H A S Q N I N V W L S  
CACAAATTACC ATCACTTGC C ATGCCAGTCA GAACATTAAAT GTTGGTTAA 100  
+1 W Y Q Q K P G D I P K L L I Y K  
GCTGGTATCA GCAGAAACCA GGAGATATCC CTAAACTATT GATCTATAAG 150  
+1 A S N L H T G V P S R F S G S S  
GCTTCCAACT TGCACACAGG CGTCCCATCA AGGTTTAGTG GCAGTGGATC 200  
+1 G T G F T L V I S S L Q P E D I A  
TGGAACAGGT TTCACATTAG TCATCAGCAG CCTGCAGCCT GAAGACATTG 250  
+1 T Y Y C Q Q G R S Y P L T F G A  
CCACTTACTA CTGT CAACAG GGTCCGAAGTT ATCCTCTCAC GTTCGGTGCT 300  
+1 G T K L E L K  
GGGACCAAGC TGGAGCTGAA A 350

Fig. 7

+1 E V Q L L E Q S G A E L V K P G A  
 GAGGTGCAGC TGCTCGAGCA GTCTGGAGCT GAGCTGGTGA AGCCTGGGGC 50  
 +1 S V K I S C K A S G Y A F S T S W  
 CTCAGTGAAG ATTTCCCTGCA AGGCTTCTTGG CTACGCCATTC AGTACCTCCT 100  
 +1 M N W V K Q R P G K G L E W I G  
 GGATGAAC TG GGTGAAACAG AGGCCTGGAA AGGCTCTTGA GTGGATTGGA 150  
 +1 R I Y P G D G D T N Y N G K F K G  
 CGGATTATC CTGGAGATGG AGATACTAAC TACAATGGGA AGTTCAAGGG 200  
 +1 K A T L T A D K S S S T A Y M Q L  
 CAAGGCCACA CTGACTGCAG ACAATCCTC CAGCACAGCC TACATGCAAC 250  
 +1 N S L T S E D S A V Y F C V R E  
 TCAACAGCCT GACATCTGAG GACTCTGCCG TCTACTTCTG TGTAAGA GAG 300  
 +1 D A Y Y S N P Y S L D Y W G Q G T  
 GATGCCCTATT ATAGTAACCC CTATAGTTTG GACTACTGGG GTCAAGGAAC 350  
 +1 S V T V S S  
 CTCAGTCACC GTCTCCTCA 400

Fig. 6



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Fig. 9

